No cases of activation of <u>Ds</u> by <u>st</u> were found in 1,818 kernels, where 1/16 are observable for concurrent losses of <u>C Sh Wx</u> on <u>st</u> kernels. Thus <u>st</u> does not carry <u>Ac</u>, and its effect of increasing stickiness in chromosomes does not result in massive activation. Whether this may be significant to the concept of <u>Ds</u> as consisting of modified heterochromatin is difficult to judge, but should be considered, in view of the "sticky" property of heterochromatin.

On one ear of the control cross, unusually strong st expression (pitted kernels) was found. Here, there were numerous non-concurrent losses of C and Wx, demonstrating clearly that st can result in endosperm mosaics.

7. High-amylose factors.

The <u>ha</u> gene reported by Kramer, and the <u>ha</u>m gene (to be designated <u>ha</u>2) both interact strongly with <u>wx</u>, giving highly collapsed kernels, variably translucent, very similar to <u>bt</u>1 kernels. The effect of <u>ha</u>1 is much greater than that of <u>ha</u>2; in fact <u>ha</u>1 <u>wx</u> kernels have a small emount of blue-staining starch, while <u>ha</u>2 <u>wx</u> kernels do not. This property of <u>ha</u>1 <u>wx</u> is similar to that reported for <u>ae wx</u> (News Letter <u>26</u>: 5, 1952). The phenotypic effect of <u>ha</u> <u>wx</u> combinations suggests that new <u>ha</u> factors might best be sought on a waxy background, where their effects are easily distinguished phenotypically.

Using this interaction, the linkage of $\underline{\text{ha}_1}$ with chromosome 5 has been confirmed, where $\underline{\text{wx}}$ $\underline{\text{T5-9c/Wx}}$ $\underline{\text{ha}_1}$ X $\underline{\text{wx}}$ $\underline{\text{ha}_1}$ shows very few collapsed kernels (4-5%). Linkage of $\underline{\text{ha}_2}$ with chromosome 10 is also clear. Chromosomes 1, 2, 3, 4, 5, 6, and 8 show independence ratios with waxy translocations, but $\underline{\text{wx}}$ $\underline{\text{T9-10b/Wx}}$ $\underline{\text{ha}_2}$ X $\underline{\text{wx}}$ $\underline{\text{ha}_2}$ shows the following:

	Normal	Waxy	Tarnished	Collapsed
ear l	(1)	53	(60)	·
ear 2	(5)	140	(122)	2
• •	(6)	193	(182)	3

9/384 = 2.3% crossing over, \underline{wx} - \underline{ha}_2 , across translocation. Anderson (Genetics, 1938) reports \underline{wx} - \underline{T} as 5.7 units. The \underline{Ha} \underline{Wx} and \underline{ha} \underline{Wx} kernels are difficult to distinguish with certainty.

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8. Allelism and mutability of anther-ear 6923.

The \underline{an}_{6923} mutant discussed last year has been tested and found allelic to \underline{an}_1 , as suggested (News Letter 30: 100, 1956). Since it is also allelic to \underline{bz}_2^m , which responds to \underline{Ac} (or \underline{M}), a test of the

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9. A crossover analysis of some mutable alleles of A1.

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An experiment has been conducted to determine whether crossing over immediately adjacent to a mutable locus can change its state of mutability (in the absence of a mutator factor) or can remove that unknown agent which according to the prevailing view of mutable loci, is suppressing the dominant allele to give the mutable recessive effect. Only those crossovers that occur immediately adjacent to the mutable locus can be expected to give the answer to this question. It is possible to make such a test using the Al locus because of its compound nature and because of the closeness of the sh2 marker. The experiment was designed as follows: Plants of α a sh/a^m Sh, dt, ac were crossed by a^S sh, Dt Dt, ac pollen and the ears produced were examined for recombinants and unusual seed types. The following diagram shows the possible types of crossovers:

	Pairing t	ype		Reg	Crossover type
A.	<u>a la 2</u>	sh		1 2	<u>α</u> <u>a^m Sh</u> α a Sh
	am	Sh		2 1	e ^m sh a sh
В.	$\frac{1}{\alpha}$ 1 a 2	sh	;	1 :	<u>a - Sh</u> <u>a a Sh</u>
	a ^m	Sh		2	a ^m sh a ^m a sh

Four different mutable alleles and one stable allele were used. Two were \underline{Dt} responding, $\underline{a^m-1}:\underline{Cache}$ and $\underline{a^m-1}:\underline{D5}$ and one was stable, $\underline{a^s}$. These have had a complete test. The other two were \underline{Ac} responding alleles, $\underline{a^{m-3}}$ and $\underline{a^{m-4}}$. They have had only a preliminary test. The \underline{a} in the $\underline{\alpha}$ \underline{a} \underline{sh} segment is the standard dottable \underline{a} found by Emerson.

The data listed in table 1 indicate that the \underline{Dt} responding alleles pair with either $\underline{\alpha}$ or \underline{a} as expected. The occurrence of $\underline{\alpha}$ \underline{a}^m \underline{Sh} cases proves pairing type A while the $\underline{\alpha}$ - \underline{Sh} and \underline{a}^m \underline{a} \underline{sh} cases prove type B. The only unusual crossover type found was a single colored non-shrunken case which had normal \underline{A} phenotype and proved to have brown pericarp. It is not known, however, whether this pericarp is dominant or recessive. If it turns out to be dominant this would be an $\underline{\alpha}$ \underline{A} \underline{Sh} case or the type expected from a crossover removal of an agent suppressing \underline{A} phenotype.